

L-Ascorbic Acid Increases NF κ B Binding Activity in UVA-Irradiated HaCaT Keratinocytes

To the Editor:

The biologic effects of UV on skin may occur either as an acute alteration shortly after UV exposure (inflammation, sunburn cells) or as a chronic damage (photoaging, photocarcinogenesis) (Beissert and Granstein, 1996). Reactive oxygen species (ROS) are involved in the pathophysiologic mechanisms leading to photooxidative damage on nucleic acids, lipids, and proteins. ROS are also responsible for the induction of proinflammatory cytokines and adhesion molecules (Punnonen *et al*, 1991; Krutmann and Grewe, 1995). ROS are constantly generated in epidermal keratinocytes and are rapidly removed by nonenzymic and enzymic antioxidant substances that prevent harmful effects of free radicals and maintain a prooxidant/antioxidant balance, thus resulting in cell and tissue stabilization (Fuchs *et al*, 1989; Yohn *et al*, 1991).

The intracellular pathway of ROS signal transduction is not fully understood. The nuclear transcription factor kappa B (NF κ B) may be one candidate for the intracellular response of UVA-induced ROS in keratinocytes. NF κ B is an inducible transcription factor involved in the regulation of genes participating in either inflammatory or immune responses. In most cell types NF κ B resides in the cytoplasm in a latent form composed of several subunits. NF κ B activation occurs by dissociation from its inhibitor protein (I κ B) and its translocation into the nucleus (Baldwin, 1996). NF κ B can be activated by various agents including UV. The fact that NF κ B activation may be inhibited by antioxidants, suggests that ROS are involved in activating NF κ B (Devary *et al*, 1993; Bender *et al*, 1998).

Activation of NF κ B can be induced in human skin by UVB (Fisher *et al*, 1996). In cultured human keratinocytes NF κ B activation occurs after combined UVA/UVB irradiation and also in cultured fibroblasts UVA irradiation activates NF κ B; however, high UVA doses lead to decreased NF κ B (Vile *et al*, 1995; Djavaheri-Mergny *et al*, 1999; Saliou *et al*, 1999). UVA-induced upregulation of interleukin-1 α (IL-1 α) can be inhibited by L-ascorbic acid (Tebbe *et al*, 1997). In this investigation, we tested the influence of L-ascorbic acid on NF κ B binding activity, in order to obtain a linkage between UVA-induced ROS and immune response in keratinocytes.

HaCaT cells, an immortalized aneuploid human keratinocyte cell line (Boukamp *et al*, 1988) shows a higher constitutive NF κ B binding activity than normal keratinocytes. The response to some cell cycle regulating factors in normal keratinocytes and HaCaT cells has been compared (Chaturvedi *et al*, 1999; Qin *et al*, 1999). A strong activation of NF κ B by factors inducing cell cycle arrest is only observed in normal keratinocytes. The absence on responsiveness to cell cycle regulatory factors correlates well with the immortalized state of HaCaT cells. In spite of the constitutive level of binding activity present in HaCaT, these cells provide a homogenous system that allow to compare other NF κ B inducing agents. Saliou *et al* (1999) showed in HaCaT cells that NF κ B

activation after UV radiation can be modulated by different agents. These studies are not possible to do by using a highly heterogeneous system like human normal keratinocytes coming from different individuals.

UVA irradiation increases TBARS formation in HaCaT keratinocytes. This effect can be partially inhibited by L-ascorbic acid indicating that ROS are involved (data not shown). NF κ B binding activity, measured in the nuclear protein extract of HaCaT keratinocytes, is only increased within a short time period after UVA irradiation (**Fig 1**). L-ascorbic acid supplementation and UVA irradiation (1 J per cm²) showed a synergistic effect on NF κ B binding activity in HaCaT keratinocytes. NF κ B binding activity was found to be increased to 156% (SEM \pm 19%) in UVA-irradiated cells. L-ascorbic acid alone did not change NF κ B binding activity; however, UVA irradiation-increased NF κ B binding activity in L-ascorbic acid supplemented cells showed remarkable increase (**Fig 2a, b**). Both NF κ B subunits p50 and p65 were detectable in nuclear protein extracts of HaCaT keratinocytes (**Fig 2c**).

The regulation of NF κ B by ROS is cell type specific. In certain cell types, such as Wurzburg subclone of T cells, L6 skeletal muscle myotubes, human breast CF-7, and 70Z/3 pre-B cells, H₂O₂ was shown to be an effective inducer of NF κ B-binding activity (Sen and Packer, 1996; Manna *et al*, 1998). It has been reported that H₂O₂ can also induce NF κ B binding activity in HeLa cells and Jurkat cells (Schreck *et al*, 1991; Brennan and O'Neill, 1995);

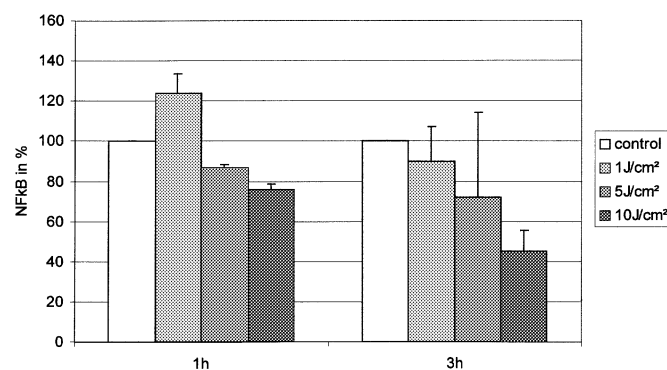


Figure 1. Highest NF κ B binding activity in HaCaT cells was seen 1 h after UVA irradiation (1 J per cm²). HaCaT cells were irradiated with 1, 5, or 10 J per cm². UVA irradiation was performed from the bottom of the cell culture dishes using a lamp with a range of 320–400 nm (Waldmann, Villingen, Germany). The output measured by a radiometer through culture dish was 6.7 mW per cm². Nuclear proteins were extracted from the cultured cells according to Corsini *et al* (1997). Electrophoretic mobility shift assays (EMSA) were performed according to Hirano *et al* (1998). A double-stranded oligonucleotide containing the NF κ B binding site (sense 5'-AGCTTCAGAGGGGACTTTCCGAG-AGG-3', antisense 5'-TCGACCTCTCGAAAGTCCCCTCTGA-3') was labeled with [α -³²P]dCTP (NEN Life Science Products, Boston, MA) using Klenow Fragment (Gibco BRL, Gaithersburg, MD). Data for NF κ B binding activity were quantified using computer-based densitometry (TINA).

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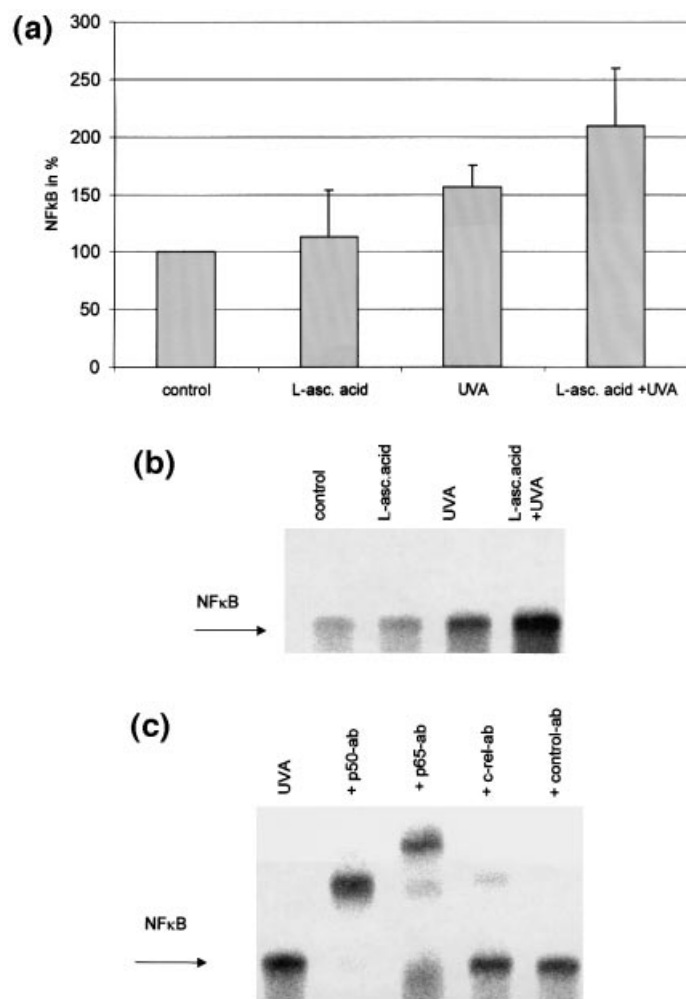


Figure 2. (a, b) L-ascorbic acid increases UVA-induced NFκB binding activity in HaCaT keratinocytes. NFκB binding activity was measured by EMSA in UVA-irradiated (1 J per cm^2) HaCaT keratinocytes supplemented with L-ascorbic acid (10^{-4} M). NFκB binding activity was quantified using computer-based densitometry and related to nonirradiated HaCaT keratinocytes without L-ascorbic acid supplementation. UVA-induced NFκB binding activity was markedly increased by L-ascorbic acid (mean \pm SEM). **(c) NFκB subunits p50 and p65 are both expressed in HaCaT cells after UVA irradiation.** NFκB supershift assay for detection of p50, p65, and c-rel subunits was used in HaCaT keratinocytes irradiated with UVA (1 J per cm^2). Both p50 and p65 specific antibodies were found positive, whereas c-rel was not detectable. Specific antibodies directed against NFκB subunits p50, p65, c-rel (Santa Cruz Biotechnology, Santa Cruz) were used for supershift assay.

however, some other groups could not detect NFκB binding activity in these cell lines after H_2O_2 stimulation (Anderson *et al*, 1994; Li and Karin, 1999).

The effect of ascorbic acid on NFκB activation varies in different cell types. Ascorbate increases the binding of NFκB to DNA in TNF- α stimulated Jurkat cells. The ability of ascorbate to enhance cytoplasmic inhibitory IκB α protein degradation correlates completely with its capacity to induce NFκB binding to DNA that potentiate NFκB-mediated transactivation of the HIV-1 long-terminal repeat (Muñoz *et al*, 1997).

Beside its antioxidative properties, ascorbic acid can also interact with metal ions that contribute to oxidative damage through the production of hydroxyl and lipid alkoxyl radicals (Carr and Frei, 1999). Ascorbate increases NO production by increasing the amount of iNOS in activated macrophages (Mizutani and Tsukagoshi, 1999). This is caused by a delay of IκB recovery and

prolonged NFκB DNA binding in lipopolysaccharide and interferon- γ activated macrophages leading to an increase in the iNOS protein level (Mizutani and Tsukagoshi, 1999).

HaCaT keratinocytes possess very efficient systems to maintain high levels of both intracellular and extracellular ascorbic acid (Savini *et al*, 1999). The regeneration and uptake of ascorbic acid from extracellular medium contributes to the intracellular antioxidant capacity, and consequently cells become more resistant to free radical generation and cell death induced by UVB irradiation (Savini *et al*, 1999).

Our results indicate that UVA induces NFκB binding activity in HaCaT keratinocytes, and this effect can be increased by L-ascorbic acid indicating that the latter has either an effect on IκB degradation or prooxidative properties. Redox regulation of cell nutrients may induce changes in cell signalling or protein conformation that represent an important mechanism in the regulation of transcription factors and gene expression.

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The Pro162 Variant is a Loss-of-Function Mutation of the Human Melanocortin 1 Receptor Gene

To the Editor:

α melanocyte-stimulating hormone (α MSH) stimulates mammalian melanocytes differentiation and/or proliferation (Abdel-Malek *et al*, 1995), by binding to the G protein-coupled melanocortin 1 receptor (MC1R) and triggering the cAMP cascade. The gene encoding for human MC1R was simultaneously cloned by Mountjoy *et al* (1992) and Chhajlani and Wikberg (1992), who reported slightly different sequences. Mammalian pigmentation depends, among other factors, on the relative amounts of skin eumelanin and pheomelanin (Thody and Graham, 1998). In man, MC1R loss-of-function mutations, such as Arg151Cys, are causally associated with predominantly pheomelanin red hair (reviewed by Rees, 2000). Because red hair is a risk factor for increased UV radiation skin sensitivity, a relationship of MC1R allelic variants and skin cancer susceptibility has been investigated. Evidence has been presented for an association between the Arg151Cys, Arg160Trp, and Asp294His variants, red hair, and melanoma (Palmer *et al*, 2000).

The study of MC1R structure–function relationships and its association with complex traits is complicated by the high polymorphism of the gene (Rees, 2000). The functional effects of many amino acid substitutions are debated or unknown. Moreover, the reported wild-type sequences differ in potentially relevant residues. An intriguing variant, originally reported as a wild-type allele, displays a proline at position 162 (Mountjoy *et al*, 1992), in the interface between the second cytosolic loop and the fourth transmembrane fragment, instead of an arginine residue reported by others (Chhajlani and Wikberg, 1992; Tan *et al*, 1999). Because Pro residues distort helical structures, an Arg/Pro substitution at the beginning of a transmembrane fragment could be functionally important.

In a survey of MC1R variants in cultured human melanoma and giant congenital nevus cells, performed by sequencing the entire coding sequence, we failed to find the Pro162 allele in 10 samples (Fig 1A, consensus sequence deposited in GenBank, number AF326275). Unfortunately, no data are available as to the pigmentation phenotypes of the patients/donors. This prompted us to screen a larger number of samples by allele-specific PCR (AS-PCR). Thirty healthy individuals were selected, of which 13 had fair skin and poor tanning ability, whereas 17 had black hair and a high basal pigmentation and tanning response. We used an artificial

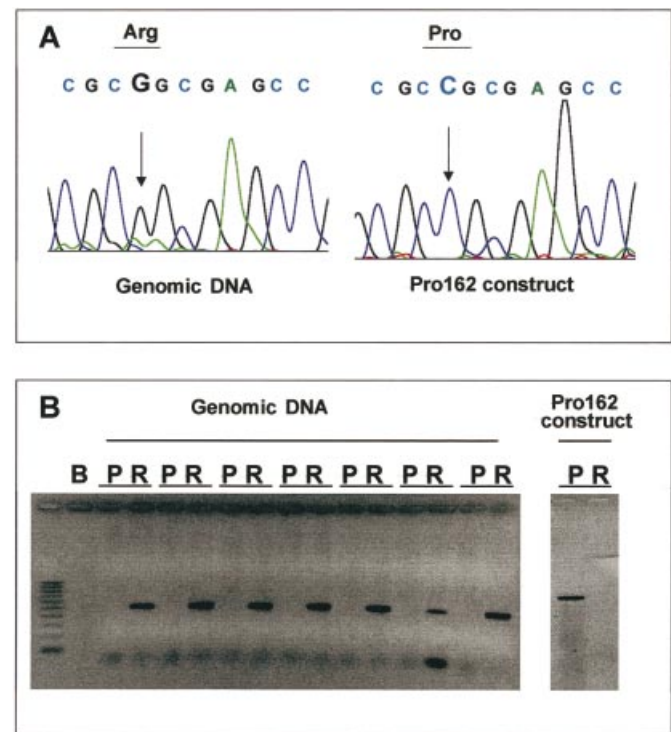


Figure 1. Identification of the Arg162 allele as the wild-type form of MC1R. The MC1R coding sequence was amplified from genomic DNA using *Pfu* polymerase and primers CCTAAGCTTACTCC-TTCCTGCTTCCTGGACA (forward, nt 435–456) and CTG-GAATTCACACTTAAAGCGCGTGCACCGC (reverse, nt 1418–1439), designed from the sequence reported by Mountjoy *et al* (1992), GenBank entry X65634, and containing added *Hind* III and *Eco* R I sites (underlined) for cloning in pcDNA3 (Invitrogen, Carlsbad, CA). (A) Sequence of amplicons from a human melanoma cell line (left), and a Pro162 construct (right), used as sequencing quality control. The relevant codon is highlighted, and the mutated base shown by an arrow. (B) Inability to detect the Pro162 variant by AS-PCR of genomic DNA. Forward primers TGACCCTGCCGCGGGCGCG (R, specific for the Arg162 form) and TGACCCTGCCGCGGGCGCC (P, for the Pro162 variant), where the nucleotide responsible for specificity is shown in bold, were used with the reverse primer described above, and 1 μ g of genomic DNA or 12 pg of the Pro162 pcDNA3 construct as target. PCR reactions with the specific primer sets for Pro (P) and Arg (R) alleles, as indicated over each lane, were analyzed by agarose gel electrophoresis. A blank reaction, without added target (labeled B), and a 100 bp ladder (first lane, left) are shown.

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